RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10 511,632
Source: PCT
Date Processed by STIC: 10-26-04

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        PATENT APPLICATION:  US/10/511,632      TIME: 17:36:19

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     3 <110> APPLICANT: Taisho Pharmaceutical Co., Ltd.
     4       IKEDA, Akiko
     5       SHINONAGA, Hideki
     6       FUJIMOTO, Natsuko
     7       KASAI, Yoko
     9 <120> TITLE OF INVENTION: HAIR GROWTH TONIC
    11 <130> FILE REFERENCE: Q84294
C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/511,632
C--> 13 <141> CURRENT FILING DATE: 2004-10-18
    13 <150> PRIOR APPLICATION NUMBER: PCT/JP03/04884
    14 <151> PRIOR FILING DATE: 2003-04-17
    16 <150> PRIOR APPLICATION NUMBER: JP 2002-115529
    17 <151> PRIOR FILING DATE: 2002-04-17
    19 <160> NUMBER OF SEQ ID NOS: 61
    21 <170> SOFTWARE: PatentIn version 3.3
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    24 <211> LENGTH: 365
    25 <212> TYPE: PRT
    26 <213> ORGANISM: HOMO SAPIENS
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    43         50                  55                  60
    46 Lys Lys Leu Cys His Leu Tyr Gln Asp His Met Gln Tyr Ile Gly Glu
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    50 Gly Ala Lys Thr Gly Ile Lys Glu Cys Gln Tyr Gln Phe Arg His Arg
    51                 85                  90                  95
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    55             100                 105                 110
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    59             115                 120                 125
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    70 Asp Trp Leu Trp Gly Gly Cys Gly Asp Asn Ile Asp Tyr Gly Tyr Arg
    71                 165                 170                 175
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79           195              200              205
82 Glu Ala Gly Arg Arg Thr Val Tyr Asn Leu Ala Asp Val Ala Cys Lys
83     210              215              220
86 Cys His Gly Val Ser Gly Ser Cys Ser Leu Lys Thr Cys Trp Leu Gln
87 225              230              235              240
90 Leu Ala Asp Phe Arg Lys Val Gly Asp Ala Leu Lys Glu Lys Tyr Asp
91           245              250              255
94 Ser Ala Ala Ala Met Arg Leu Asn Ser Arg Gly Lys Leu Val Gln Val
95           260              265              270
98 Asn Ser Arg Phe Asn Ser Pro Thr Thr Gln Asp Leu Val Tyr Ile Asp
99     275              280              285
102 Pro Ser Pro Asp Tyr Cys Val Arg Asn Glu Ser Thr Gly Ser Leu Gly
103     290              295              300
106 Thr Gln Gly Arg Leu Cys Asn Lys Thr Ser Glu Gly Met Asp Gly Cys
107 305              310              315              320
110 Glu Leu Met Cys Cys Gly Arg Gly Tyr Asp Gln Phe Lys Thr Val Gln
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138 ggcccaggtt ggggagagga cggagggtgg ccgcagcggg ttcctgagtg aattacccag  360
140 gagggactga gcacagcacc aactagagag gggtcagggg gtgcgggact cgagcgagca  420
142 ggaaggaggc agcgcctggc accagggctt tgactcaaca gaattgagac acgtttgtaa  480
144 tcgctggcgt gccccgcgca caggatccca gcgaaaatca gatttcctgg tgaggttgcg  540
146 tgggtggatt aatttggaaa aagaaactgc ctatatcttg ccatcaaaaa actcacggag  600
148 gagaagcgca gtcaatcaac agtaaactta agagacccc  gatgctcccc tggtttaact  660
150 tgtatgcttg aaaattatct gagagggaat aaacatcttt tccttcttcc ctctccagaa  720
152 gtccattgga atattaagcc caggagttgc tttggggatg gctggaagtg caatgtcttc  780
154 caagttcttc ctagtggctt tggccatatt tttctccttc gcccaggttg taattgaagc  840
156 caattcttgg tggtcgctag gtatgaataa ccctgttcag atgtcagaag tatatattat  900
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160 ccacttgtat caggaccaca tgcagtacat cggagaaggc gcgaagacag gcatcaaaga 1020
162 atgccagtat caattccgac atcgacggtg gaactgcagc actgtggata acacctctgt 1080
164 ttttggcagg gtgatgcaga taggcagccg cgagacggcc ttcacatacg ccgtgagcgc 1140
166 agcaggggtg gtgaacgcca tgagccgggc gtgccgcgag ggcgagctgt ccacctgcgg 1200
168 ctgcagccgc gccgcgcgcc ccaaggacct gccgcgggac tggctctggg gcggctgcgg 1260
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174 cgaggccggc cgcaggacgg tgtacaacct ggctgatgtg gcctgcaagt gccatggggt 1440
176 gtccggctca tgtagcctga agacatgctg gctgcagctg cagacttcc gcaaggtggg 1500
178 tgatgccctg aaggagaagt acgacagcgc ggcggccatg cggctcaaca gccggggcaa 1560
180 gttggtacag gtcaacagcc gcttcaactc gcccaccaca caagacctgg tctacatcga 1620
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184 cctgtgcaac aagacgtcgg agggcatgga tggctgcgag ctcatgtgct gcggccgtgg 1740
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188 ctacgtcaag tgcaagaagt gcacggagat cgtggaccag tttgtgtgca agtagtgggt 1860
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192 tggttttttgg tttttagaaa tatttttttat ttttccccaa gaattgcaac cggaaccatt 1980
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198 tacaagactt cttttggata gtatagaatg aaggggggaa taacacatac cctaacttag 2160
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208 cattttcaaa atgataattt acaatggaag gacaagaatg tcatattctc aaggaaaaaa 2460
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260 ctcttaattt attgcataat gatatccaca tcagccaact gtggctcttt aatttattgc 4020
262 ataatgatat tcacatcccc tcagttgcag tgaattgtga gcaaaagatc ttgaaagcaa 4080
264 aaagcactaa ttagtttaaa atgtcacttt tttggttttt attatacaaa aaccatgaag 4140
266 tactttttt atttgctaaa tcagattgtt cctttttagt gactcatgtt tatgaagaga 4200
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272 gtcttgcgtg atttgtatat ttcactggtt taaaaaacaa acatcgaaag gcttattcca   4380
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278 <211> LENGTH: 2460
279 <212> TYPE: DNA
280 <213> ORGANISM: HOMO SAPIENS
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287 ttaccctggg gttgacagtc tcgccagagg tctcatttca tactgtcttt tcggatctga    180
289 tcctcttggt aaacaggcgg ggatgtttta ccctacagag ccgatgtatg tgtgagttcg    240
291 ctgtgagttc tttgagtgtc tcaaacttgt ggggcctttt ctcggttgca ctgggattga    300
293 agagggaaga ggcccaaggt gtttccgggc aagcggcggg gttaagtgga gatgcgactc    360
295 gtgaggctct cctttccgat ccccctttgg gacaccctct gcctacctct accctggagc    420
297 cagggagacc caagtcttgg tgaccggatg ggcccgctct cagttggcct gggctctggg    480
299 aactggtgga ctctccctgg gggcttcggg ctgggagtgg gttcggtttg tgtggcttcg    540
301 gctctaacaa agagatccgc tgtaatccgc cgaatctgtt atcaatttct ctgctgcttg    600
303 agccccgccc cacgcgcccc gcccgccgcg aagcttggaa agtgcacgcg gccagcacca    660
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307 tgtaaatgtg cacgagtgtg aatgtgtatg atgtgtgtgc acgcggcatc ggctgccctt    780
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315 cttggggaca actggcctgg tgttggagcc ctgcttagca ggcgctgggg accacataag   1020
317 cattcctctt tggagaagcc ccgaagcgtc caggccaaag ggggcggttc acggaagaaa   1080
319 aaccttgcac gcccttgagc gcatagcttt accagggctg cctaggtccc gcctcttgcc   1140
321 cttttacggc acaggttcca agccaggctc ttcccaccgc cttaaagagg ctcacctttc   1200
323 ttttcttttc tgtggaaggg gctccttcag gggctatggg cgatgcagtg cggcagggtt   1260
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335 aaccaagtct gagttgggct ggagaggcta gactggaggc agggtggcag agttccaacg   1620
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355 acccctgcc cttccctccc gcgtcctgcc cccatcctcc acccccgcg ctggccaccc    2220
357 cgcctccttg gcagcctctg gcggcagcgc gctccactcg cctcccgtgc tcctctcgcc   2280
359 catggaatta attctggctc cacttgttgc tcggcccagg ttggtgagag gacggagggt   2340
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381 Cys Gln Glu Ile Thr Val Pro Met Cys Arg Gly Ile Gly Tyr Asn Leu
382   35   40   45
385 Thr His Met Pro Asn Gln Phe Asn His Asp Thr Gln Asp Glu Ala Gly
386   50   55   60
389 Leu Glu Val His Gln Phe Trp Pro Leu Val Glu Ile Gln Cys Ser Pro
390 65   70   75   80
393 Asp Leu Arg Phe Phe Leu Cys Thr Met Tyr Thr Pro Ile Cys Leu Pro
394   85   90   95
397 Asp Tyr His Lys Pro Leu Pro Pro Cys Arg Ser Val Cys Glu Arg Ala
398   100   105   110
401 Lys Ala Gly Cys Ser Pro Leu Met Arg Gln Tyr Gly Phe Ala Trp Pro
402   115   120   125
405 Glu Arg Met Ser Cys Asp Arg Leu Pro Val Leu Gly Arg Asp Ala Glu
406   130   135   140
409 Val Leu Cys Met Asp Tyr Asn Arg Ser Glu Ala Thr Thr Ala Pro Pro
410 145   150   155   160
413 Arg Pro Phe Pro Ala Lys Pro Thr Leu Pro Gly Pro Pro Gly Ala Pro
414   165   170   175
417 Ala Ser Gly Gly Glu Cys Pro Ala Gly Gly Pro Phe Val Cys Lys Cys
418   180   185   190
421 Arg Glu Pro Phe Val Pro Ile Leu Lys Glu Ser His Pro Leu Tyr Asn
422   195   200   205
425 Lys Val Arg Thr Gly Gln Val Pro Asn Cys Ala Val Pro Cys Tyr Gln
426   210   215   220
429 Pro Ser Phe Ser Ala Asp Glu Arg Thr Phe Ala Thr Phe Trp Ile Gly
430 225   230   235   240
433 Leu Trp Ser Val Leu Cys Phe Ile Ser Thr Ser Thr Thr Val Ala Thr
434   245   250   255
437 Phe Leu Ile Asp Met Asp Thr Phe Arg Tyr Pro Glu Arg Pro Ile Ile
438   260   265   270
441 Phe Leu Ser Ala Cys Tyr Leu Cys Val Ser Leu Gly Phe Leu Val Arg
442   275   280   285
445 Leu Val Val Gly His Ala Ser Val Ala Cys Ser Arg Glu His Asn His
446   290   295   300
449 Ile His Tyr Glu Thr Thr Gly Pro Ala Leu Cys Thr Ile Val Phe Leu
450 305   310   315   320
453 Leu Val Tyr Phe Phe Gly Met Ala Ser Ser Ile Trp Trp Val Ile Leu
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457 Ser Leu Thr Trp Phe Leu Ala Ala Ala Met Lys Trp Gly Asn Glu Ala

VERIFICATION SUMMARY   DATE: 10/26/2004
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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date